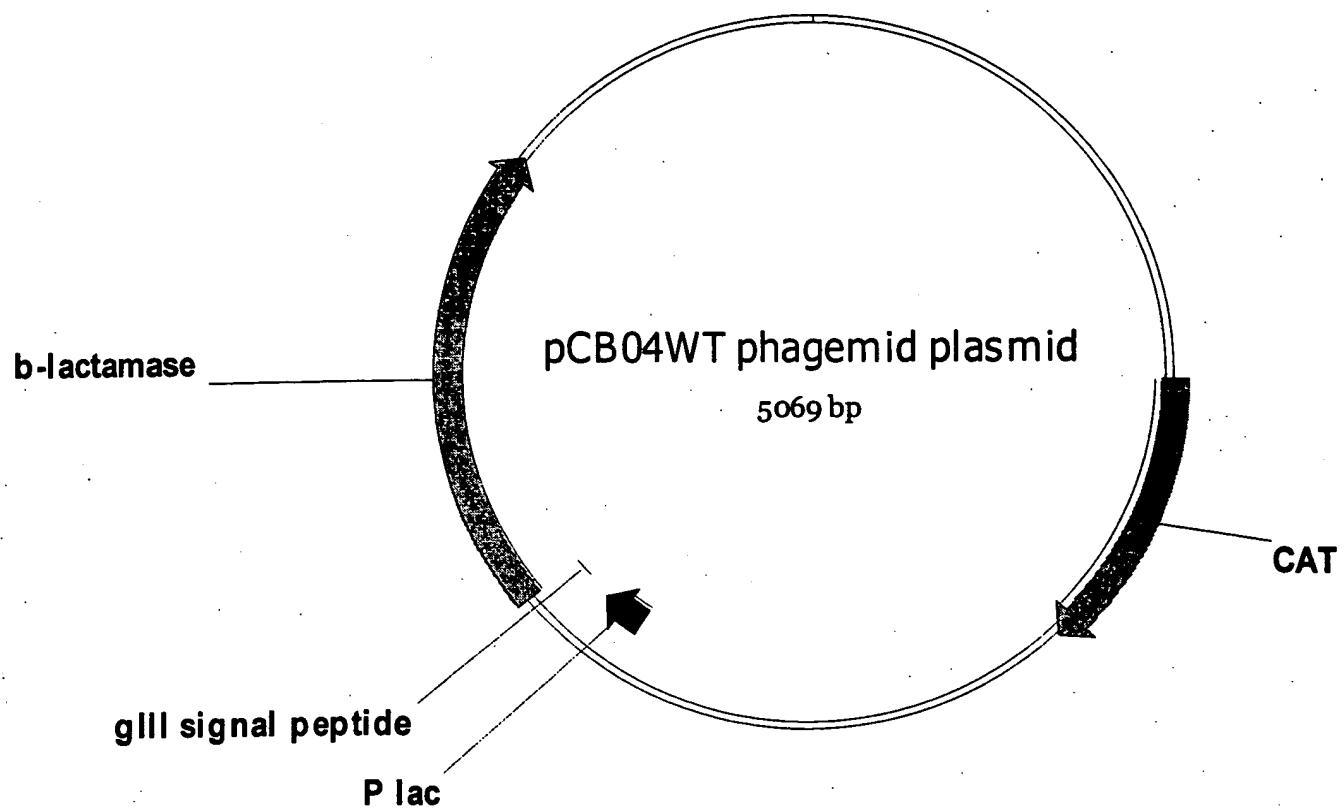


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FIGURE 1.



## FIGURE 2

FIGURE 3.

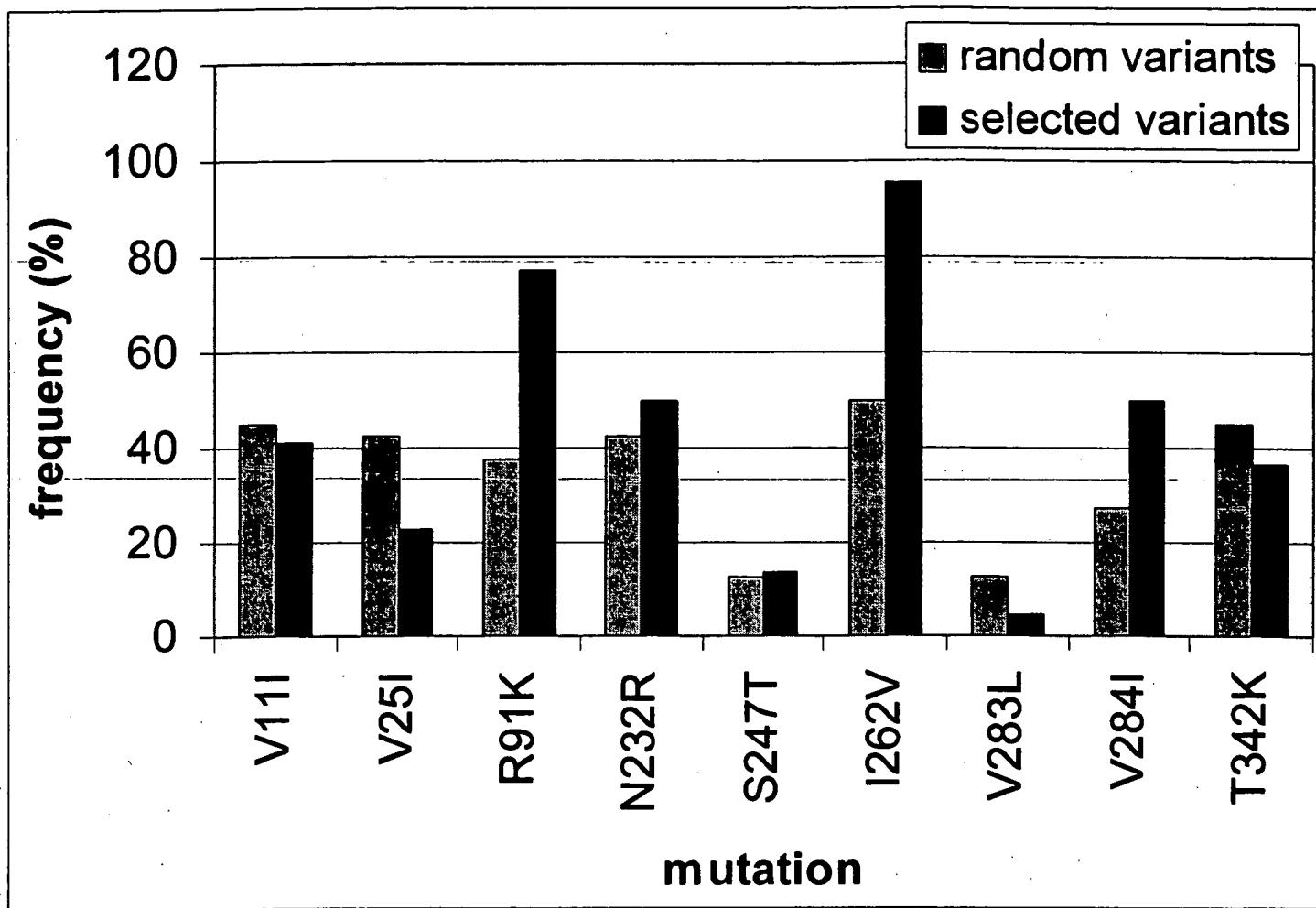
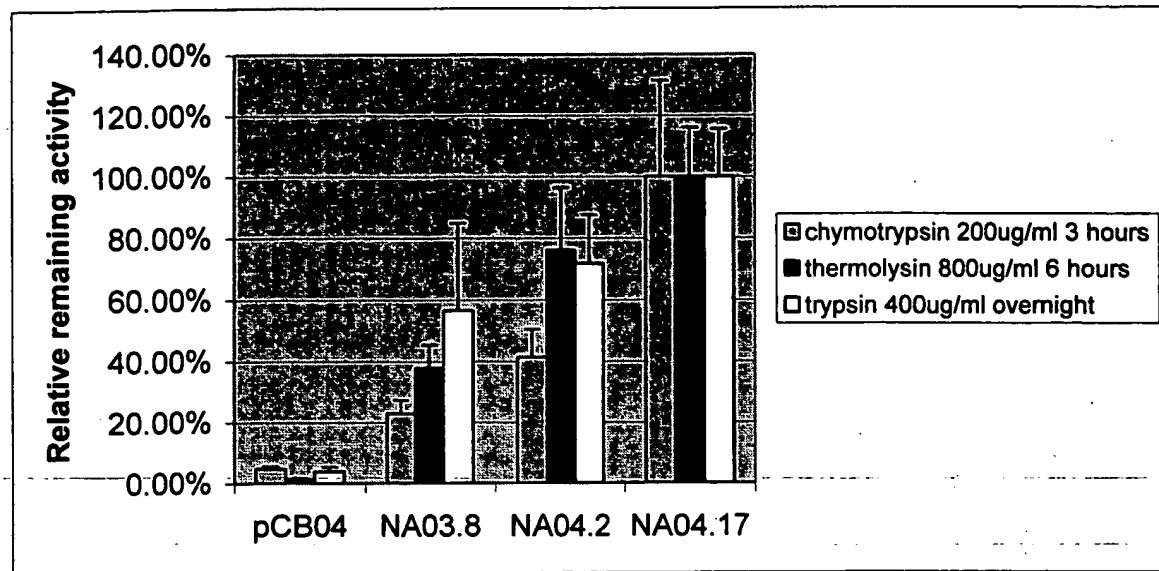


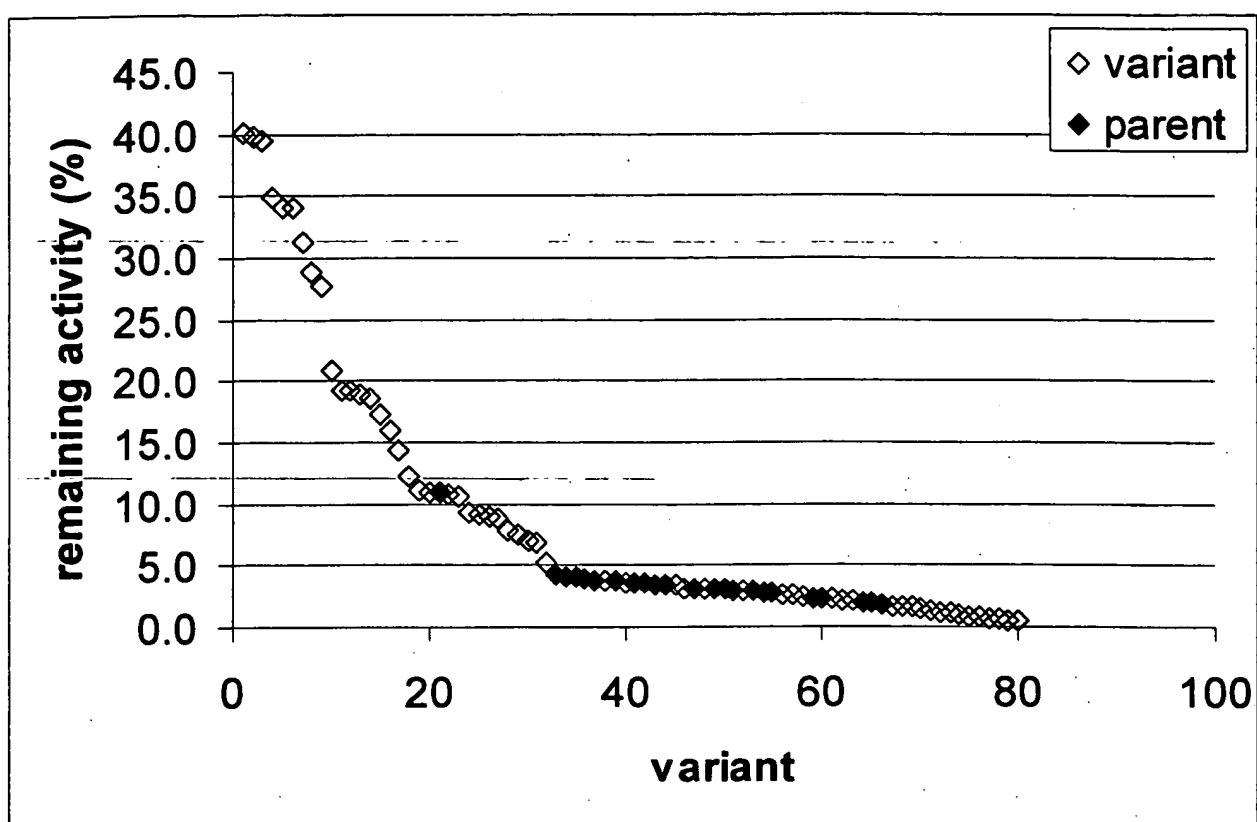
FIGURE 4.

**FIGURE 5.**



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FIGURE 6.



**FIGURE 7.**

heavy chain:  
QvkIqqsgaelvrsqtsvklsctasgfmikdsymhwIrqgpeeqglewigwidpengdteyapkfqgkattfttdtsntaylqlisslttsedavyycnegtpgpyyfdywgqgtvt  
vss (SEQ ID NO:2)

linker:

Gggsgggsggggs (SEQ ID NO:3)

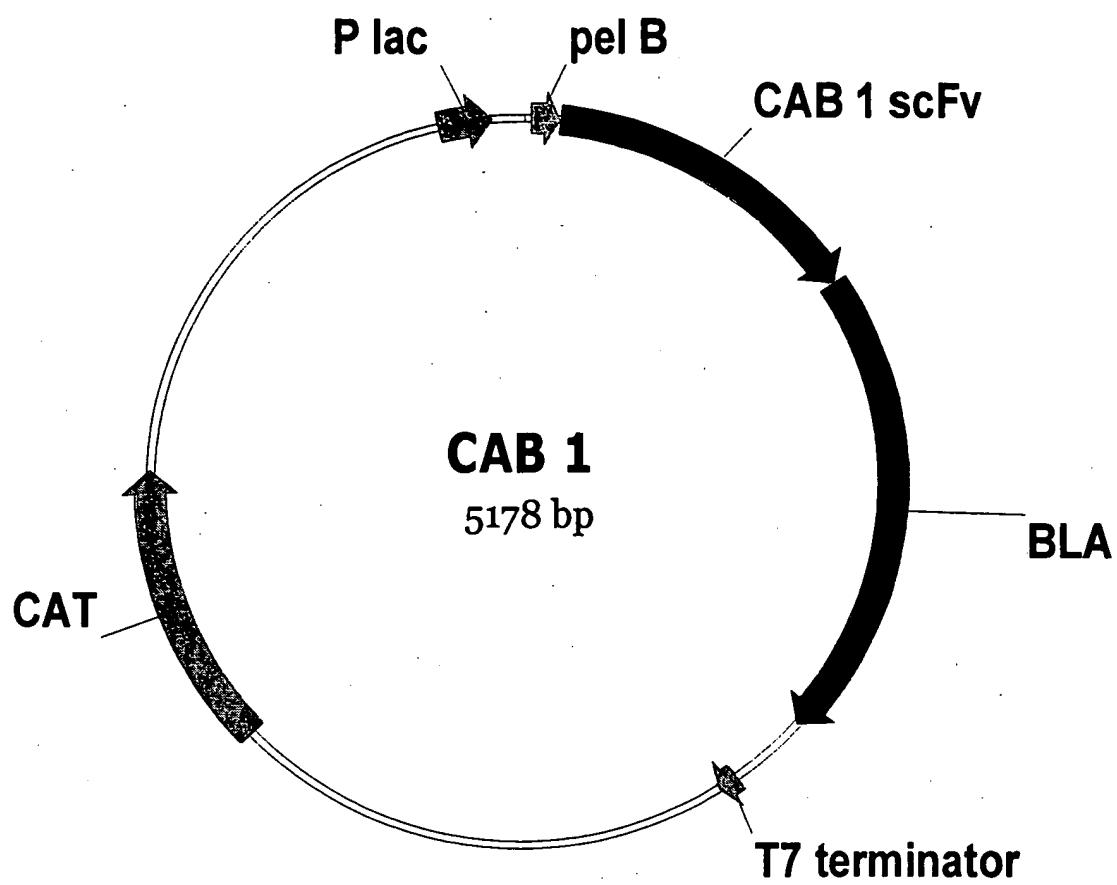
light chain:

Envltqspaimsaspgekvititcsassssvsymhwffqqkpgtspklwystslnlasgvparfsgsgstsylsmeaedaatyyccqrssyplifgagtklelkraat (SEQ ID NO:4)

BLA:

Lvsekqlaevvantitplmkadqsypgmaavaviyqgkphyytfgkadiaankpvtptqtlfelgsiskftfgvlgddaiargeislddavtrywpqltgkqwqgirmldlatyttaggpl  
qypdevtdnasllrfyqnwqpqkwkpgttlyanasiglsgalavkpsgmpyeqamitrvlpkldhtwinvpkaeeahyawgyrdgkavrvspgmldaqaygvktnvqdm  
anwvmanmapenvadasllkgqialaqsrwrigsmyqglgwemlnwpeantvetsfgnwalaplvpaevnppappkaswhktgstggfsyvafipekqigivml  
antsypparveaayhilealq (SEQ ID NO:5)

**FIGURE 8.**



## **FIGURE 9.**

FIGURE 10.

Translation of CAB1  
human consensus

1 (1) QVKLQOSGAEVLRSGTSVKLSCTASGFNIKDSYMMHWLRQGPQEGLWIGWIDPENGTEYAPKFQGKATTFITDTSSNTAY (SEQ ID NO:6)  
(1) Q KP G TFS V A G RF I R K

80

Translation of CAB1  
human consensus

81 (81) LQLSSLTSED TAVYYCNEGTPGYYFDYWGQGTIVTSSGGGGSGGGSGGGSENVLTQSPAIMSASPGEKVTITCSA (SEQ ID NO:7)  
(81) PR QS P S V QT

160

Translation of CAB1  
human consensus

161 (161) SSSVSYMHHWFQQKPGTSPKIWIYSTNLASGVPARFSGSGSTSYSLTISRMEAEDAATYYCQQRSSYPLTFGAGTKLELKRAA (SEQ ID NO:8)  
(161) Y V D K A L E D G

240

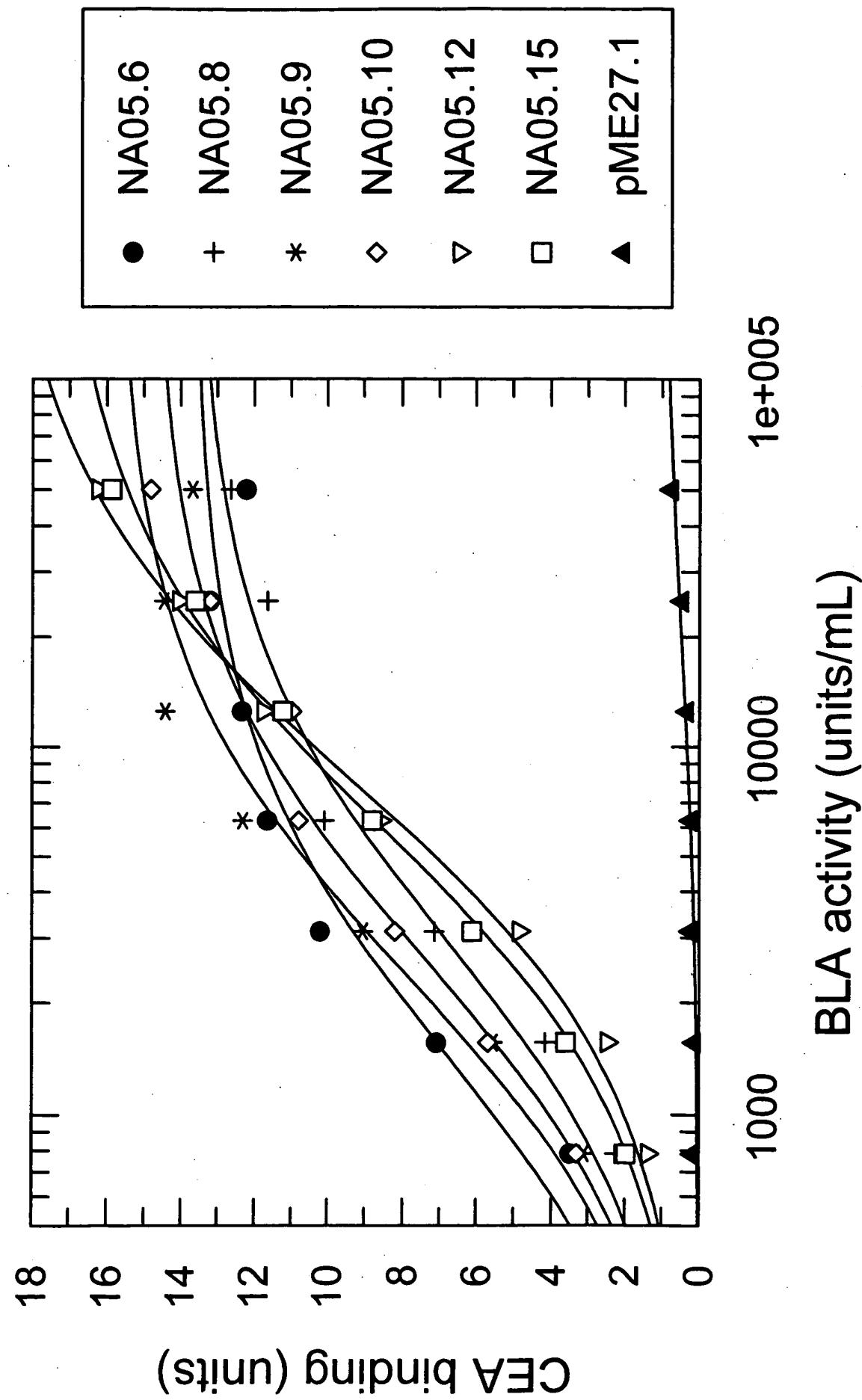


FIGURE 12.

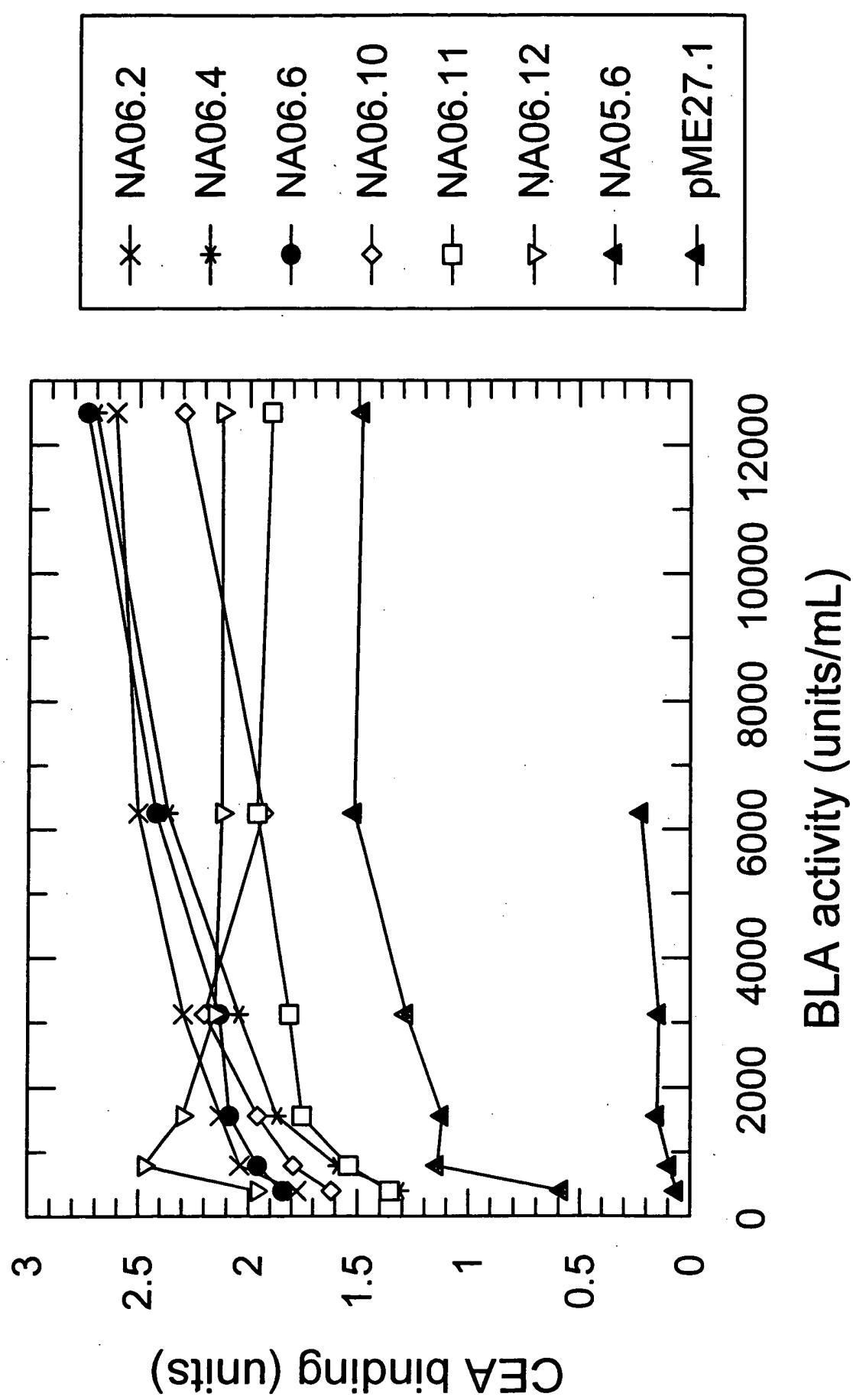


FIGURE 13.

